

## FIGURE 1

	Antibody				
Native	Signal S	Sequer	ce and	Variable	Region

Nucleotide Sequence SEQ ID NO:1 Amino Acid Sequence SEQ ID NO:2

			_		_	-							
ATG Met 1	GAG Glu	ACA Thr	GAC Asp	ACA Thr 5	ATC Ile	CTG Leu	CTA Leu	TGG Trp	GTG Val 10	CTG Leu	CTG Leu	CTC Leu	39
TGG Trp	GTT Val 15	CCA Pro	GGC Gly	TCC Ser	ACT Thr	GGT Gly 20	GAC Asp	ATT Ile	GTG Val	CTG Leu	ACC Thr 25	CAA Gln	78
TCT Ser	CCA Pro	GCT Ala	TCT Ser	TTG Leu	GCT Ala	GTG Val	TCT Ser	CTA Leu 35	GGG Gly	CAG Gln	AGG Arg	GCC Ala	117
ACC Thr 40	ATC Ile	TCC Ser	TGC <b>C</b> ys	AAG Lys	GCC Ala 45	AGC Ser	CAA Gln	AGT Ser	GTT Val	GAT Asp 50	TAT Tyr	GAT Asp	156
GGT Gly	GAT Asp	AGT Ser 55	TAT Tyr	ATG Met	AAC Asn	TGG Trp	TAC Tyr 60	CAA Gln	CAG Gln	AAA Lys	CCA Pro	GGA Gly 65	195
CAG Gln	CCA Pro	CCC Pro	AAA Lys	CTC Leu 70	CTC Leu	ATC Ile	TAT Tyr	GCT Ala	GCA Ala 75	TCC Ser	AAT Asn	CTA Leu	234
GAA Glu	TCT Ser 80	GGG Gly	ATC Ile	CCA Pro	GCC Ala	AGG Arg 85	TTT Phe	AGT Ser	GGC Gly	AGT Ser	GGG Gly 90	TCT Ser	273
GGG Gly	ACA Thr	GAC Asp	TTC Phe 95	ACC Thr	CTC Leu	AAC Asn	ATC Ile	CAT His 100	CCT Pro	GTG Val	GAG Glu	GAG Glu	312
GAG Glu 105	GAT Asp	GCT Ala	GCA Ala	ACC Thr	TAT Tyr 110	TAC Tyr	TGT Cys	CAG Gln	CAA Gln	AGT Ser 115	AAT Asn	GAG Glu	351
GAT <u>Asp</u>	CCT Pro	CCG Pro 120	ACG Thr	TTC Phe	GGT Gly	GGA Gly	GGC Gly 125	ACC Thr	AAG Lys	CTG Leu	GAA Glu	ATC Ile 130	390
AAA Lys													396

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## Figure 2

Murine Antibody 3B9 Heavy Chain Native Signal Sequence and Variable Region

Nucleotide Sequence SEQ ID NO: 3 Amino Acid Sequence SEQ ID NO: 4

GAATTCGCGG CCGCTATGCA GGGACAATCA GCAGCAGCAA	40													
TGAGGAAGTA AGCCTGTGCA GAT ATG AAC AGG CTT ACT TCC Met Asn Arg Leu Thr Ser  1 5  TCA TTG CTG CTG CTG ATT GTC CCT GCA TAT GTC CTC TCC														
TCA TTG CTG CTG CTG ATT GTC CCT GCA TAT GTC CTG TCC Ser Leu Leu Leu Ile Val Pro Ala Tyr Val Leu Ser 10 15	120													
CAG GTT ACT CTG AAA GAG TCT GGC CCT GGG ATA TTG CAG Gln Val Thr Leu Lys Glu Ser Gly Pro Gly Ile Leu Gln 20 25 30	159													
CCC TCC CAG ACC CTC AGT CTG ACT TGT TCT TTC TCT GGG Pro Ser Gln Thr Leu Ser Leu Thr Cys Ser Phe Ser Gly 35 40 45	198													
TTT TCA CTG AGC ACT TCT GGT ATG GGT GTG AGC TGG ATT Phe Ser Leu Ser Thr Ser Gly Met Gly Val Ser Trp Ile 50 55	237													
CGT CAG CCT TCA GGA AAG GGT CTG GAG TGG CTG GCA CAC Arg Gln Pro Ser Gly Lys Gly Leu Glu Trp Leu Ala His 60 65 70	276													
ATT TAC TGG GAT GAT GAC AAG CGC TAT AAC CCA TCC CTG  Ile Tyr Trp Asp Asp Asp Lys Arg Tyr Asn Pro Ser Leu  75 80	315													
AAG AGC CGG CTC ACA ATC TCC AAG GAT ACC TCC AGC AAC  Lys Ser Arg Leu Thr Ile Ser Lys Asp Thr Ser Ser Asn  85 90 95	354													
CAG GTA TTC CTC AAG ATC ACC AGT GTG GAC ACT GCA GAT Gln Val Phe Leu Lys Ile Thr Ser Val Asp Thr Ala Asp 100 105 110	393													
ACT GCC ACA TAC TAC TGT GCT CGA AGA GAG ACT GTG TTC Thr Ala Thr Tyr Tyr Cys Ala Arg Arg Glu Thr Val Phe 115	432													

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# Figure 2 (con't)

TAC Tyr	TGG Trp 125	TAC Tyr	TTC Phe	GAT Asp	GTC Val	TGG Trp 130	GGC Gly	GCA Ala	GGG Gly	ACC Thr	ACG Thr 135	GTC Val	471
		TCC Ser											483

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#### FIGURE 3

Human/Murine 3B9 Chimeric Antibody Heavy Chain Signal Sequence and Variable Region

Nucleotide Sequence SEQ ID NO: 9 Amino Acid Sequence SEQ ID NO:10

AMINO AC	id sequence	SEQ ID N	0:10	
	TTG CAG ACC Leu Gln Thr 5			
	TCT GGT GCC Ser Gly Ala			Glu
	CCT GGG ATA Pro Gly Ile 30			
	TGT TCT TTC Cys Ser Phe			
	GGT GTG AGC Gly Val Ser 55			
	GAG TGG CTG Glu Trp Leu 70			
	TAT AAC CCA Tyr Asn Pro			Ile
	GAT ACC TCC Asp Thr Ser 95			
	GTG GAC ACT Val Asp Thr			
	AGA GAG ACT Arg Glu Thr 120			
	GCA GGG ACC Ala Gly Thr 135	Thr Val	Thr Val	423

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#### FIGURE 4

Humanized 3B9 Antibody Heavy Chain Signal Sequence and Variable Region

Nucleotide Sequence SEQ ID NO: 11
Amino Acid Sequence SEO ID NO: 12

	Amir	no Ad	cid :	Seque	ence	SEQ	ID I	10:	12					
ŧ,	ATG Met 1	<i>GTG</i> Val	<i>TTG</i> Leu	<i>CAG</i> Gln	ACC Thr 5	<i>CAG</i> Gln	<i>GTC</i> Val	TTC Phe	ATT Ile	TCT Ser 10	<i>CTG</i> Leu	<i>TTG</i> Leu	<i>CTC</i> Leu	39
	TGG Trp	ATC Ile 15	TCT Ser	GGT Gly	GCC Ala	TAC Tyr	GGG Gly 20	CAG Gln	GTT Val	ACC Thr	CTG Leu	CGT Arg 25	GAA Glu	78
	TCC Ser	GGT Gly	CCG Pro	GCA Ala 30	CTA Leu	GTT Val	AAA Lys	CCG Pro	ACC Thr 35	CAG Gln	ACC Thr	CTG Leu	ACG Thr	117
	TTA Leu 40	ACC Thr	TGC Cys	ACC Thr	TTC Phe	TCC Ser 45	GGT Gly	TTC Phe	TCC Ser	CTG Leu	TCG Ser 50	ACC Thr	TCC Ser	156
	GGT Gly	ATG <u>Met</u>	GGT Gly 55	GTT Val	TCC Ser	TGG Trp	ATC Ile	CGT Arg 60	CAG Gln	CCG Pro	CCG Pro	GGT Gly	AAA Lys 65	195
	GGT Gly	CTA Leu	GAA Glu	TGG Trp	CTG Leu 70	GCT Ala	CAC <u>His</u>	ATC Ile	TAC Tyr	TGG Trp 75	GAC Asp	GAC Asp	GAC Asp	234
	AAA Lys	CGT <u>Arq</u> 80	TAC Tyr	AAC Asn	CCG Pro	AGC Ser	CTG Leu 85	AAA Lys	TCC Ser	CGT Arg	CTG Leu	ACG Thr 90	ATA Ile	273
	TCC Ser	AAA Lys	GAC Asp	ACC Thr 95	TCC Ser	CGT Arg	AAC Asn	CAG Gln	GTT Val 100	GTT Val	CTG Leu	ACC Thr	ATG Met	312
	ACT Thr 105	AAC Asn	ATG Met	GAC Asp	CCG Pro	GTT Val 110	GAC Asp	ACC Thr	GCT Ala	ACC Thr	TAC Tyr 115	TAC Tyr	TGC Cys	351
	GCT Ala	CGA Arg	CGC <u>Arg</u> 120	GAA Glu	Thr	GTT Val	TTC Phe	TAC Tyr 125	TGG Trp	TAC Tyr	TTC Phe	GAC Asp	GTT <u>Val</u> 130	390
•	TGG Trp	GGT Gly	CGT Arg	GGT Gly	ACC Thr 135	CCA Pro	GTT Val	ACC Thr	GTG Val	AGC Ser 140	TCA Ser			423

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## FIGURE 5

Humania	zed	3B9	Αn	tibo	ody	Light	Chain
Signal	Seq	uenc	e	and	Var	iable	Region

Nucleotide Sequence SEQ ID NO: 13 Amino Acid Sequence SEQ ID NO: 14

			Jeque	SIICE	SEQ	10 1	10:	L 4					
ATG Met 1	GGA Gly	TGG Trp	AGC Ser	TGT Cys 5	ATC Ile	ATC Ile	CTC Leu	TTC Phe	TTG Leu 10	GTA Val	GCA Ala	ACA Thr	39
GCT Ala	ACA Thr 15	GGT Gly	GTC Val	CAC His	TCC Ser	GAT Asp 20	ATC Ile	GTG Val	ATG Met	ACC Thr	CAG Gln 25	TCT Ser	78
CCA Pro	GAC Asp	TCG Ser	CTA Leu 30	GCT Ala	GTG Val	TCT Ser	CTG Leu	GGC Gly 35	GAG Glu	AGG Arg	GCC Ala	ACC Thr	<b>117</b>
ATC Ile 40	AAC Asn	TGC Cys	AAG Lys	GCC Ala	TCC Ser 45	CAA Gln	AGT Ser	GTT Val	GAT Asp	TAT Tyr 50	GAT Asp	GGT Gly	156
GAT <u>Asp</u>	AGT Ser	TAT Tyr 55	Met	AAC Asn	TGG Trp	TAT Tyr	CAG Gln 60	CAG Gln	AAA Lys	CCC Pro	GGG Gly	CAG Gln 65	195
CCT Pro	CCT Pro	AAG Lys	TTG Leu	CTC Leu	Ile	TAC Tyr	GCT Ala	GCA Ala	TCC Ser	AAT Asn	CTA Leu	GAA Glu	234

							GAA Glu	
		70			75			

TCT	GGG	GTA	CCT	GAC	CGA	TTC	AGT	GGC	AGC	GGG	TCT	GGG	273
<u> 261</u>	GIY	Val	PLO	ASP	Arg	Pne	Ser	GIY	ser	GIA	Ser	Gly	
	80					85					90	_	

ACA.	GAT	TTC	ACT	CTC	ACC	ATC	AGC	AGC	CTG	CAG	GCT	GAA	312
Thr	Asp	Phe	Thr	Leu	Thr	Ile	Ser	Ser	Leu	Gln	Ala	Glu	
			95					100					

GAT	GTG	GCA	GTA	TAC	TAC	TGT	CAG	CAA	AGT	AAT	GAG	GAT Asp	351
r2b	AGI	VIG.	AGI	ryr	ryr	Cys	GIL	Gin	ser	<u>ASN</u>	Glu	ASP	
105					110	_				115			

							AAA Lys	390
	120	-	•	125	•		130	

CGT	393
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#### FIGURE 6A

Signa:	l Sequ	ience	2	
Nucle	otide	SEQ	ID	NO:5
Amino	Acid	SEQ	ID	NO: 6

ATG GTG TTG CAG ACC CAG GTC TTC ATT TCT CTG TTG CTC
Met Val Leu Gln Thr Gln Val Phe Ile Ser Leu Leu Leu

1 10

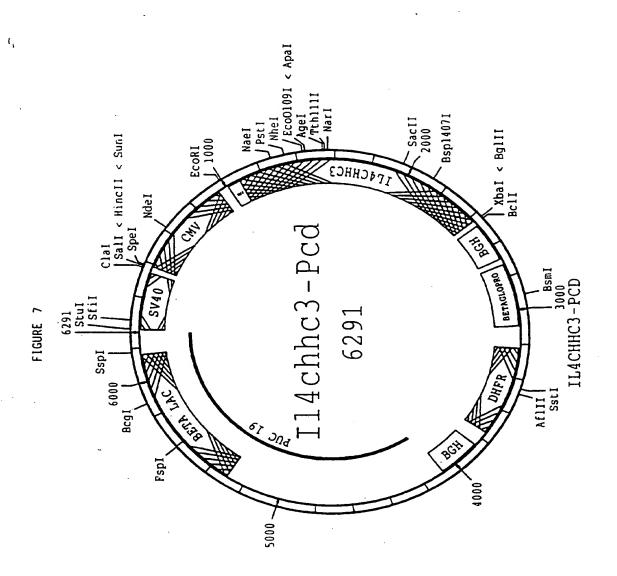
TGG ATC TCT GGT GCC TAC Trp Ile Ser Gly Ala Tyr 15

#### FIGURE 6B

Signal Sequence Nucleotide SEQ ID NO:7 Amino Acid SEQ ID NO:8

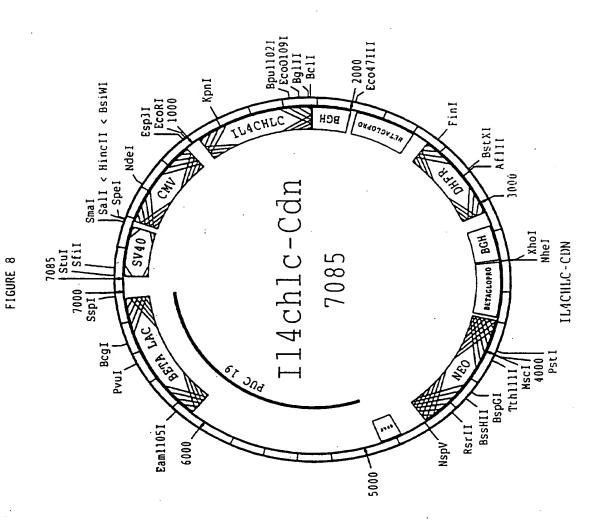
ATG GGA TGG AGC TGT ATC ATC CTC TTC TTG GTA GCA ACA
Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr
1 5 10

GCT ACA GGT GTC CAC TCC GAT ATC GTG ATG ACC CAG TCT 78
Ala Thr Gly Val His Ser Asp
15 20



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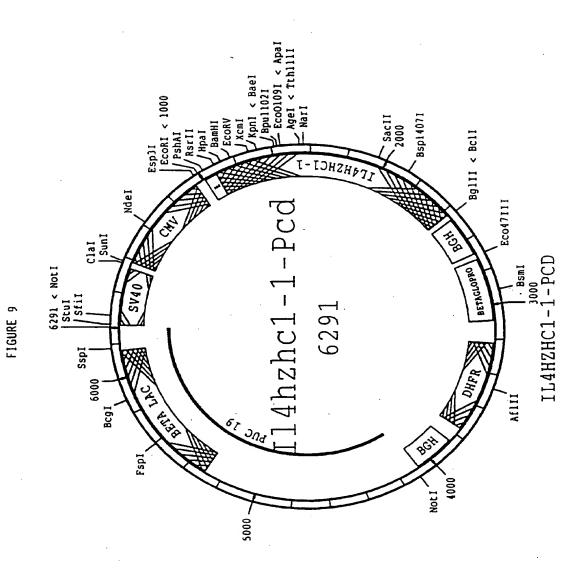
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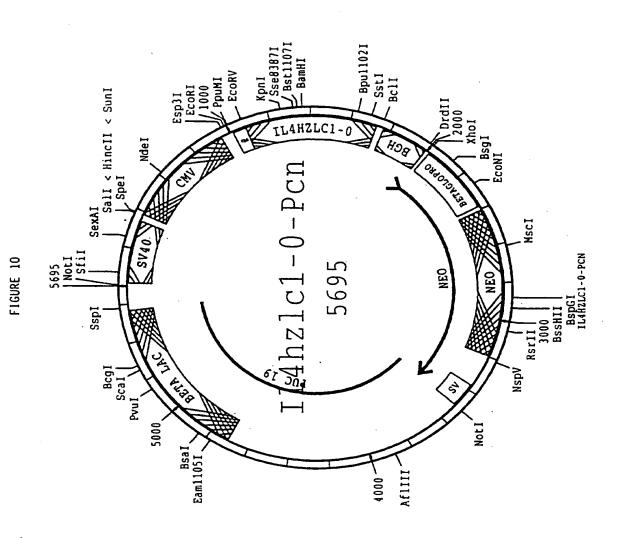




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